Nicolas Pollet

List of Publications by Year in descending order

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69 4,781 27 65
papers citations h-index g-index

74 74 74 7460
all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Evaluation of time profile reconstruction from complex two-color microarray designs. BMC Bioinformatics, 2008, 9, 1.	2.6	875
2	The Genome of the Western Clawed Frog <i>Xenopus tropicalis</i> . Science, 2010, 328, 633-636.	12.6	708
3	Synexpression groups in eukaryotes. Nature, 1999, 402, 483-487.	27.8	386
4	Comparison of T7E1 and Surveyor Mismatch Cleavage Assays to Detect Mutations Triggered by Engineered Nucleases. G3: Genes, Genomes, Genetics, 2015, 5, 407-415.	1.8	260
5	Gene expression screening in Xenopus identifies molecular pathways, predicts gene function and provides a global view of embryonic patterning. Mechanisms of Development, 1998, 77, 95-141.	1.7	198
6	Mutations in JAGGED1 gene are predominantly sporadic in Alagille syndrome. Gastroenterology, 1999, 116, 1141-1148.	1.3	178
7	Nrarp is a novel intracellular component of the Notch signaling pathway. Genes and Development, 2001, 15, 1885-1899.	5.9	153
8	The transmembrane protein XFLRT3 forms a complex with FGF receptors and promotes FGF signalling. Nature Cell Biology, 2004, 6, 38-44.	10.3	149
9	Microbiota and Mucosal Immunity in Amphibians. Frontiers in Immunology, 2015, 6, 111.	4.8	128
10	Periodic repression of Notch pathway genes governs the segmentation of Xenopus embryos. Genes and Development, 1999, 13, 1486-1499.	5.9	127
11	Xenbase: a Xenopus biology and genomics resource. Nucleic Acids Research, 2007, 36, D761-D767.	14.5	116
12	Increased XRALDH2 activity has a posteriorizing effect on the central nervous system of Xenopus embryos. Mechanisms of Development, 2001, 101, 91-103.	1.7	112
13	Minimum information specification for in situ hybridization and immunohistochemistry experiments (MISFISHIE). Nature Biotechnology, 2008, 26, 305-312.	17.5	111
14	Pax3 and Zic1 trigger the early neural crest gene regulatory network by the direct activation of multiple key neural crest specifiers. Developmental Biology, 2014, 386, 461-472.	2.0	111
15	Human and other mammalian genomes contain transposons of themarinerfamily. FEBS Letters, 1995, 368, 541-546.	2.8	81
16	Generation of trangenic Xenopus laevis using the Sleeping Beauty transposon system. Transgenic Research, 2006, 15, 751-760.	2.4	66
17	Cyclic expression of esr9 gene in Xenopus presomitic mesoderm. Differentiation, 2003, 71, 83-89.	1.9	62
18	An atlas of differential gene expression during early Xenopus embryogenesis. Mechanisms of Development, 2005, 122, 365-439.	1.7	60

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19	Compound toxicity screening and structure–activity relationship modeling in <i>Escherichia coli</i> Biotechnology and Bioengineering, 2012, 109, 846-850.	3.3	50
20	Identification of post-transcriptionally regulated Xenopus tropicalis maternal mRNAs by microarray. Nucleic Acids Research, 2006, 34, 986-995.	14.5	48
21	An ontology for Xenopus anatomy and development. BMC Developmental Biology, 2008, 8, 92.	2.1	48
22	Transgenesis procedures in <i>Xenopus</i> . Biology of the Cell, 2008, 100, 503-529.	2.0	48
23	Identification of CUG-BP1/EDEN-BP target mRNAs in Xenopus tropicalis. Nucleic Acids Research, 2008, 36, 1861-1870.	14.5	45
24	Reliability of gene expression ratios for cDNA microarrays in multiconditional experiments with a reference design. Nucleic Acids Research, 2004, 32, 29e-29.	14.5	30
25	Whispering to the Deaf: Communication by a Frog without External Vocal Sac or Tympanum in Noisy Environments. PLoS ONE, 2011, 6, e22080.	2.5	30
26	Validation of novel reference genes for RTâ€qPCR studies of gene expression in Xenopus tropicalis during embryonic and postâ€embryonic development. Developmental Dynamics, 2013, 242, 709-717.	1.8	30
27	Cloning and Characterization of Plx2 and Plx3, Two Additional Polo-like Kinases from Xenopus laevis. Experimental Cell Research, 2001, 270, 78-87.	2.6	29
28	Characterization of multiple lineages of Tc1-like elements within the genome of the amphibian Xenopus tropicalis. Gene, 2005, 349, 187-196.	2.2	29
29	Characterization of a novel <i>Xenopus tropicalis</i> cell line as a model for in vitro studies. Genesis, 2012, 50, 316-324.	1.6	28
30	How minute sooglossid frogs hear without a middle ear. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 15360-15364.	7.1	26
31	ncRNAclassifier: a tool for detection and classification of transposable element sequences in RNA hairpins. BMC Bioinformatics, 2012, 13, 246.	2.6	25
32	A large scale screen for neural stem cell markers in <i>Xenopus</i> retina. Developmental Neurobiology, 2012, 72, 491-506.	3.0	25
33	Identification of the pre–T-cell receptor α chain in nonmammalian vertebrates challenges the structure–function of the molecule. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 19991-19996.	7.1	23
34	Implication of thyroid hormone signaling in neural crest cells migration: Evidence from thyroid hormone receptor beta knockdown and NH3 antagonist studies. Molecular and Cellular Endocrinology, 2017, 439, 233-246.	3.2	23
35	Axeldb: a Xenopus laevis database focusing on gene expression. Nucleic Acids Research, 2000, 28, 139-140.	14.5	22
36	Reduced levels of survival motor neuron protein leads to aberrant motoneuron growth in a Xenopus model of muscular atrophy. Neurogenetics, 2010, 11, 27-40.	1.4	22

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37	A new tree-frog genus and species from Ivory Coast, West Africa (Amphibia: Anura: Hyperoliidae). Zootaxa, 2009, 2044, 23-45.	0.5	21
38	Construction of a 3.7-Mb Physical Map within Human Chromosome 20p12 Ordering 18 Markers in the Alagille Syndrome Locus. Genomics, 1995, 27, 467-474.	2.9	19
39	Database of queryable gene expression patterns for <i>Xenopus</i> . Developmental Dynamics, 2009, 238, 1379-1388.	1.8	19
40	The mariner Transposons Belonging to the irritans Subfamily Were Maintained in Chordate Genomes by Vertical Transmission. Journal of Molecular Evolution, 2006, 62, 53-65.	1.8	18
41	Characterization of a Xenopus tropicalis Endogenous Retrovirus with Developmental and Stress-Dependent Expression. Journal of Virology, 2011, 85, 2167-2179.	3.4	15
42	Insights on genome size evolution from a miniature inverted repeat transposon driving a satellite DNA. Molecular Phylogenetics and Evolution, 2014, 81, 1-9.	2.7	15
43	Nuclear Importation of Mariner Transposases among Eukaryotes: Motif Requirements and Homo-Protein Interactions. PLoS ONE, 2011, 6, e23693.	2.5	15
44	Exploring nervous system transcriptomes during embryogenesis and metamorphosis in Xenopus tropicalis using EST analysis. BMC Genomics, 2007, 8, 118.	2.8	14
45	Evading the annotation bottleneck: using sequence similarity to search non-sequence gene data. BMC Bioinformatics, 2008, 9, 442.	2.6	14
46	The olig family: phylogenetic analysis and early gene expression in Xenopus tropicalis. Development Genes and Evolution, 2007, 217, 485-497.	0.9	13
47	Construction of an Integrated Physical and Gene Map of Human Chromosome 20p12 Providing Candidate Genes for Alagille Syndrome. Genomics, 1997, 42, 489-498.	2.9	11
48	In situ analysis of gene expression in Xenopus embryos. Comptes Rendus - Biologies, 2003, 326, 1011-1017.	0.2	11
49	Features of the mammal mar1 transposons in the human, sheep, cow, and mouse genomes and implications for their evolution. Mammalian Genome, 2000, 11, 1111-1116.	2.2	10
50	Expression Profiling by Systematic High-Throughput In Situ Hybridization to Whole-Mount Embryos. , 2001, 175, 309-321.		10
51	Generation of BAC Transgenic Tadpoles Enabling Live Imaging of Motoneurons by Using the Urotensin II-Related Peptide (ust2b) Gene as a Driver. PLoS ONE, 2015, 10, e0117370.	2.5	10
52	Post-transcriptional regulation in Xenopus embryos: role and targets of EDEN-BP. Biochemical Society Transactions, 2005, 33, 1541.	3.4	9
53	TBid mediated activation of the mitochondrial death pathway leads to genetic ablation of the lens inXenopus laevis. Genesis, 2007, 45, 1-10.	1.6	9
54	Synchrotron analysis of a â€~mummified' salamander (Vertebrata: Caudata) from the Eocene of Quercy, France. Zoological Journal of the Linnean Society, 2016, 177, 147-164.	2.3	9

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55	A new method for long-read sequencing of animal mitochondrial genomes: application to the identification of equine mitochondrial DNA variants. BMC Genomics, 2020, 21, 785.	2.8	9
56	Tissue-specific expression of Sarcoplasmic/Endoplasmic Reticulum Calcium ATPases (ATP2A/SERCA) 1, 2, 3 during Xenopus laevis development. Gene Expression Patterns, 2011, 11, 122-128.	0.8	8
57	Deleted Chromosome 20 from a patient with Alagille syndrome isolated in a cell hybrid through leucine transport selection: study of three candidate genes. Mammalian Genome, 1994, 5, 663-669.	2.2	7
58	Ancient Adaptive Lateral Gene Transfers in the Symbiotic Opalina–Blastocystis Stramenopile Lineage. Molecular Biology and Evolution, 2020, 37, 651-659.	8.9	7
59	Evolutionary Dynamics of the Repetitive DNA in the Karyotypes of Pipa carvalhoi and Xenopus tropicalis (Anura, Pipidae). Frontiers in Genetics, 2020, 11, 637.	2.3	7
60	Gut microbial ecology of Xenopus tadpoles across life stages. , 0, 1 , .		7
61	Irradiation damage to frog inner ear during synchrotron radiation tomographic investigation. Journal of Electron Spectroscopy and Related Phenomena, 2009, 170, 37-41.	1.7	5
62	Expression of immune genes during metamorphosis of Xenopus: a survey. Frontiers in Bioscience - Landmark, 2010, 15, 348.	3.0	5
63	Databases of Gene Expression in Xenopus Development. Methods in Molecular Biology, 2012, 917, 319-345.	0.9	5
64	Properties of the various Botmar1 transcripts in imagoes of the bumble bee, Bombus terrestris (Hymenoptera: Apidae). Gene, 2007, 390, 52-66.	2.2	4
65	Identification of novel cis-regulatory elements of Eya1 in Xenopus laevis using BAC recombineering. Scientific Reports, 2017, 7, 15033.	3.3	2
66	Construction and characterization of a BAC library for functional genomics in Xenopus tropicalis. Developmental Biology, 2017, 426, 255-260.	2.0	2
67	A New Method for Sequencing the Mitochondrial Genome by Using Long Read Technology. Methods in Molecular Biology, 2021, 2277, 331-343.	0.9	1
68	Transcriptomic analysis of the trade-off between endurance and burst-performance in the frog Xenopus allofraseri. BMC Genomics, 2021, 22, 204.	2.8	1
69	Large Scale Expression Screening Identifies Molecular Pathways and Predicts Gene function. , 2002, , 27-35.		0